

IN THE SPECIFICATION

Please replace the “Detailed Description of the Drawing” with the following:

FIG. 1 presents the comparative amino acid sequence of the following members of the cystein-rich protein family of growth-regulating proteins: Cyr61 (SEQ ID NO:2); CEF10 (SEQ ID NO:20); Fisp12 (SEQ ID NO:6); CTGF (SEQ ID NO:8); and Nov (SEQ ID NO:21).

Please replace the paragraph beginning at page 23, line 13 of the specification as filed with the following:

Alternatively, human *cyr61* cDNA clones were identified with probes generated by RT-PCR. In particular, the probe for screening the human placental cDNA library was a PCR fragment generated with degenerate primers by RT-PCR of total RNA from logarithmically growing WI38 cells. The primers were derived from the sequences corresponding to the most conserved region of the open reading frame of the mouse *cyr61* cDNA. One primer, designated H61-5 [5'GGGAATTCTG(TC)GG(GATC)TG(TC)TG(TC)AA(GA)GT(GC)TG-3' (SEQ ID NO: 18)], contains a degenerate sequence, which, with the exception of the “GGGAATTC” sequence at the 5' end which was used to introduce an *Eco*RI site, is derived from nucleotides 327-346 (sense strand) of the mouse *cyr61* sequence set forth in SEQ ID NO:1. The degeneracies appear in positions corresponding to the third position of codons in SEQ ID NO:1. The second primer used for PCR amplification of a human *cyr61* sequence was designated H61-3 [5'-CCGGATCC(GA)CA(GA)TT(GA)TA(GA)TT(GA)CA-3' (SEQ ID NO: 19)], which, with the exception of the 5' sequence “CCGGATCC” used to introduce a *Bam*HI site, corresponds to the anti-sense strand complementary to nucleotides 1236-1250 of the mouse *cyr61* sequence set forth in SEQ ID NO:1. The degeneracies occur in positions complementary to the third positions of codons in mouse *cyr61* as set forth in SEQ ID NO:1. The amplified *cyr61* cDNA was cloned into the pBlueScript SK+ vector (Stratagene, La Jolla, CA) and sequenced with a Sequenase II kit (U.S. Biochemicals, Cleveland, OH).